

<b>Notice of References Cited</b>	Application/Control No. 10/530,104	Applicant(s)/Patent Under Reexamination IBBERSON ET AL.	
	Examiner Elly-Gerald Stoica	Art Unit 1647	Page 1 of 2

**U.S. PATENT DOCUMENTS**

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
*	A	US-4,383,033	05-1983	Scherberg, Neal H.	435/68.1
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

**FOREIGN PATENT DOCUMENTS**

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N	WO/94/01548	01-1994	WIPO	Sibson et al.	C12N 15/11
	O					
	P					
	Q					
	R					
	S					
	T					

**NON-PATENT DOCUMENTS**

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	Skolnick et al. From genes to protein structure and function: novel applications of computational approaches in the genomic era. Trends in Biotech 18(1): 34-39, 2000.
	V	Bork, A. Powers and pitfalls in sequence analysis: the 70% hurdle. Genome Res 10: 398-400, 2000.
	W	Doerks et al. Protein annotation: detective work for function prediction. Trends in Genetics 14(6): 248-250, 1998.
	X	Smith et al. The challenges of genome sequence annotation or "The devil is in the details". Nature Biotech 15: 1222-1223, 1997.

\*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)  
Dates in MM-YYYY format are publication dates. Classifications may be US or foreign.

<b>Notice of References Cited</b>	Application/Control No. 10/530,104	Applicant(s)/Patent Under Reexamination IBBERSON ET AL.	
	Examiner Elly-Gerald Stoica	Art Unit 1647	Page 2 of 2

**U.S. PATENT DOCUMENTS**

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-			
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

**FOREIGN PATENT DOCUMENTS**

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N					
	O					
	P					
	Q					
	R					
	S					
	T					

**NON-PATENT DOCUMENTS**

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	Brenner, S.E. Errors in genome function. Trends in Genetics 15(4): 132-133, 1999.
	V	Bork et al. Go hunting in sequence databases but watch out for the traps. Trends in Genetics. 12(10): 425-427, 1996.
	W	Wells. J.A. Additivity of mutational effects in proteins. Biochemistry 29 (37): 8509-8517, 1990.
	X	Ngo et al. Computational complexity, protein structure prediction, and the Levinthal paradox. The Protein Folding Problem and Tertiary Structure Prediction, pp. 492-495, 1994.

\*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)  
Dates in MM-YYYY format are publication dates. Classifications may be US or foreign.